



Biogeochemistry of a late marginal coccolithophorid bloom in the Bay of Biscay

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Introduction

*role of **primary production**, **calcification** and **export** processes
in **coccolithophore blooms***



the **net ecosystem dynamics**
(**primary production**,
calcification
and **pelagic respiration**)



the link between
the **bacterial community**,
TEP formation
and **carbon export**

Dynamics of coccolithophore blooms:
Contribution in climate regulation ?





Introduction

Coccolithophores:

Diatoms → Coccolithophores

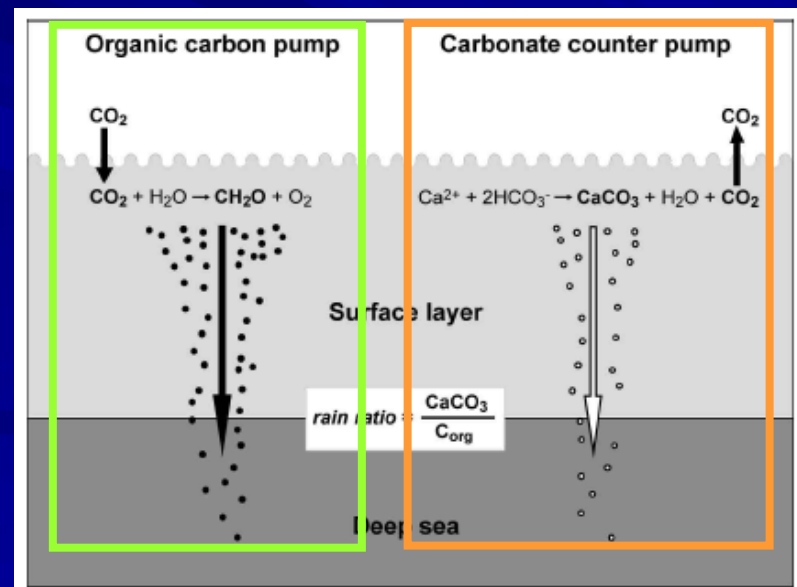
Low requirements for inorganic phosphorus and micronutrients

« Organic phase » followed by the « calcifying phase » → release of coccoliths

With regard to the C-cycle:

Organic or biological pump
(photosynthesis)

Carbonate counter-pump
(biocalcification)



From Rost & Riebesell, 2004



Introduction

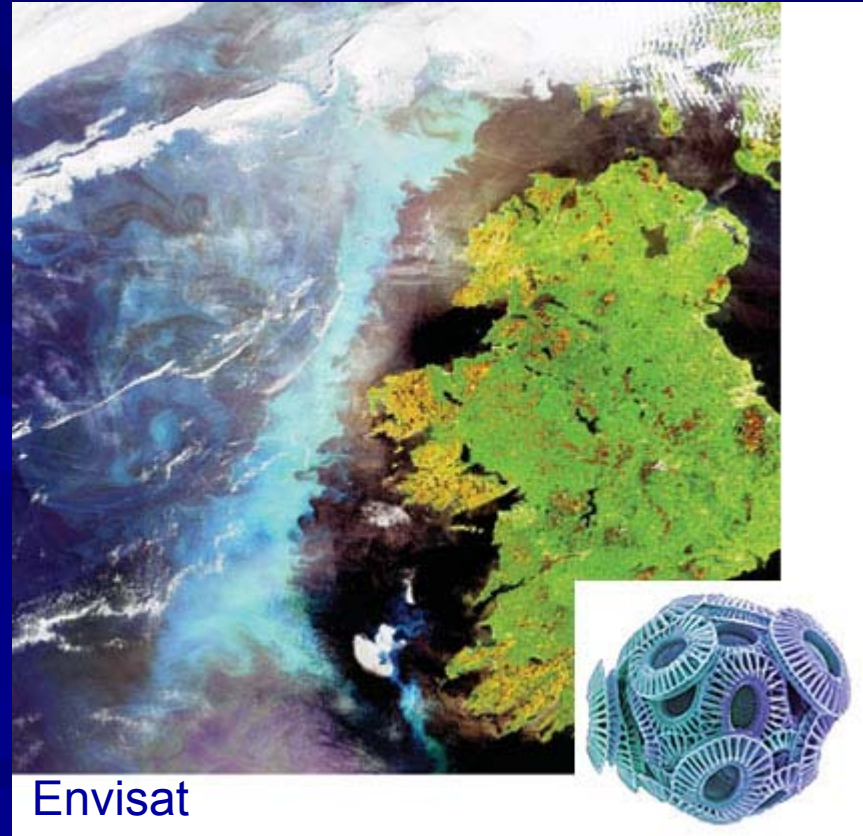


(Nature, 441, 15 June 2006)

June 2006

stretched over **500 km**

probably due to the common coccolithophore species *Emiliana huxleyi*

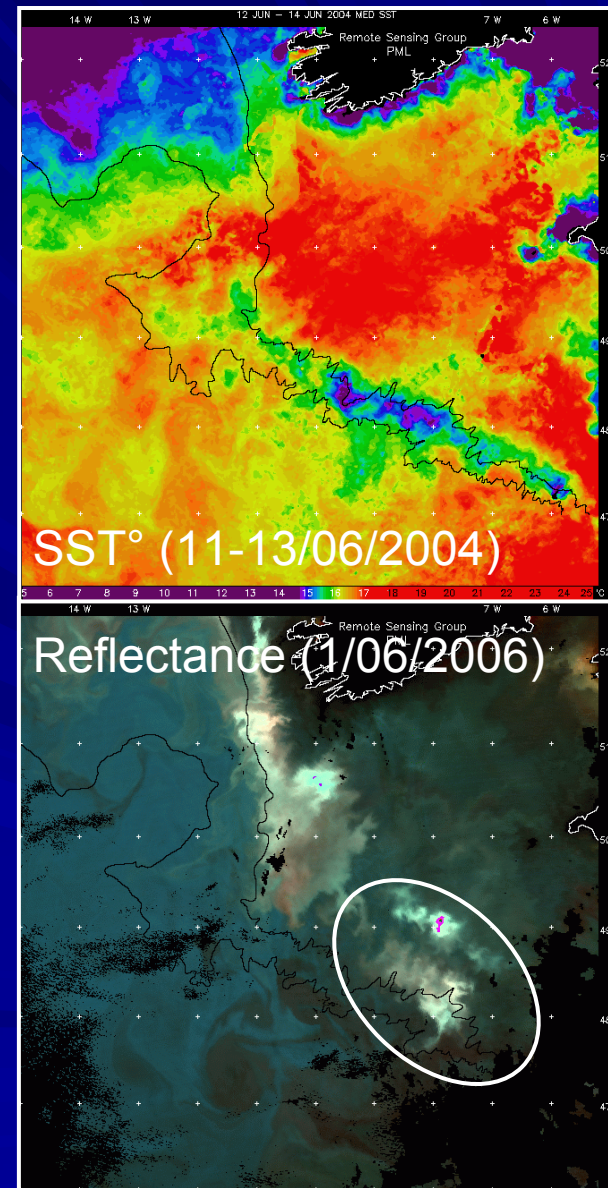
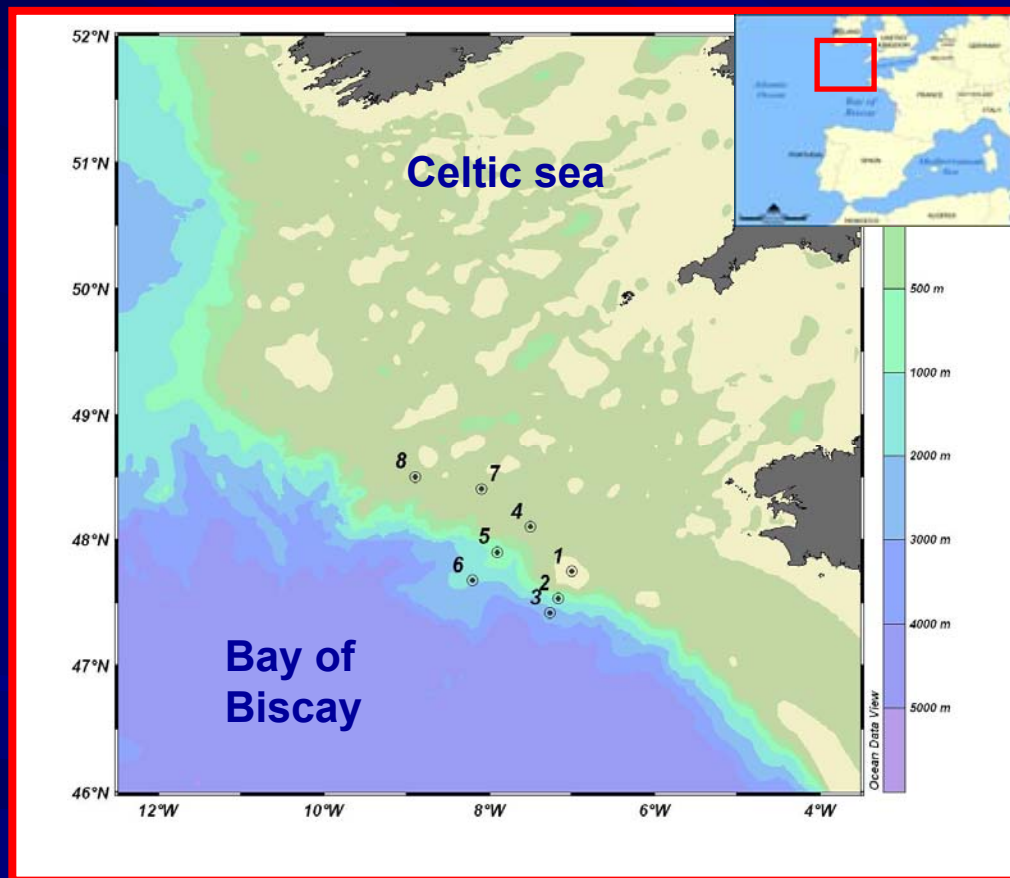


Envisat



Zone of study

31th May – 10th June 2006





Zone of study

T° : 13-15 °C

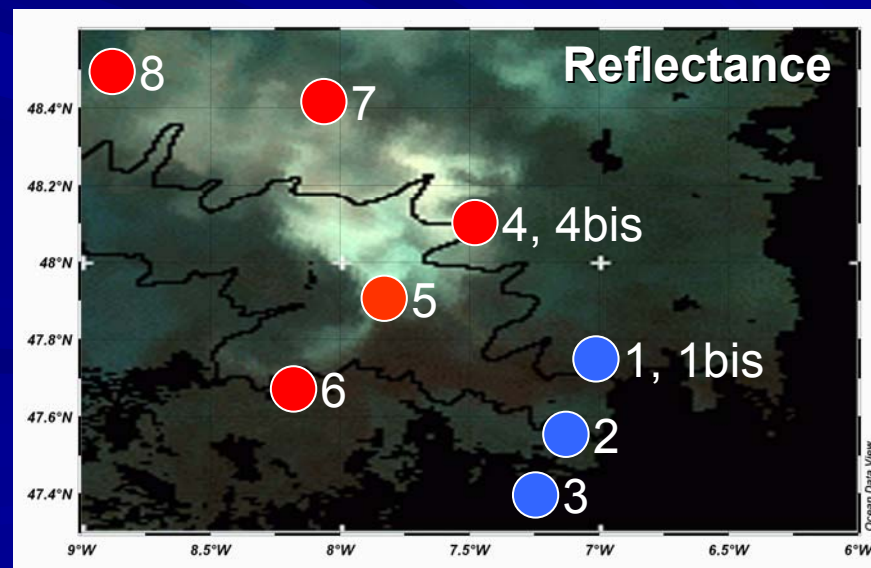
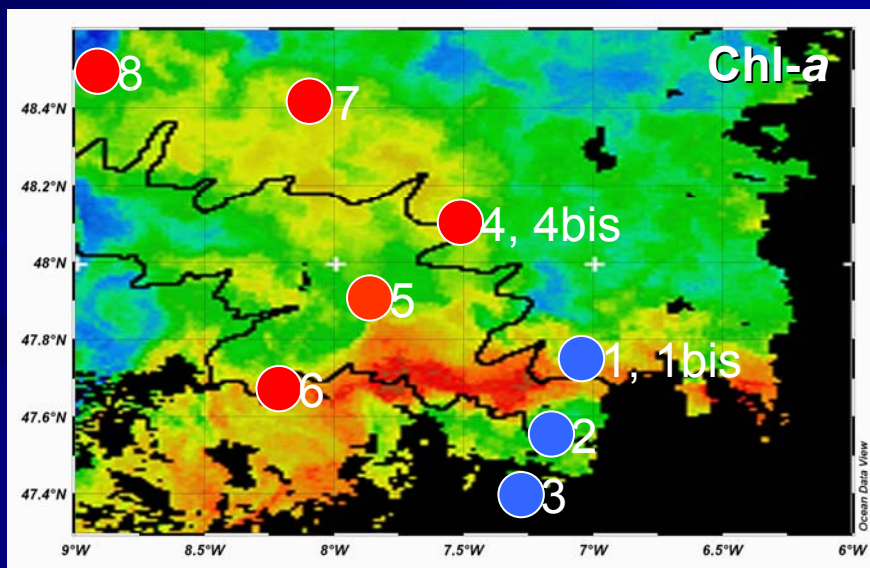
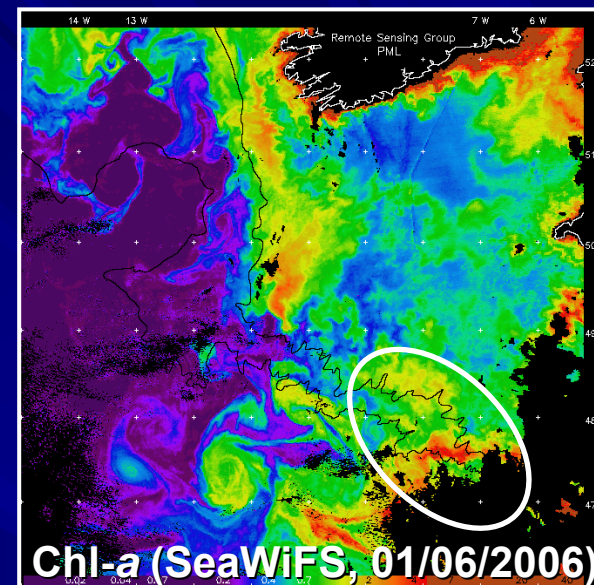
Thermocline 40-50 m

Sal: 35.5 – 35.7

$[PO_4] \sim 0 \mu M$

$[Si] < 2 \mu M$

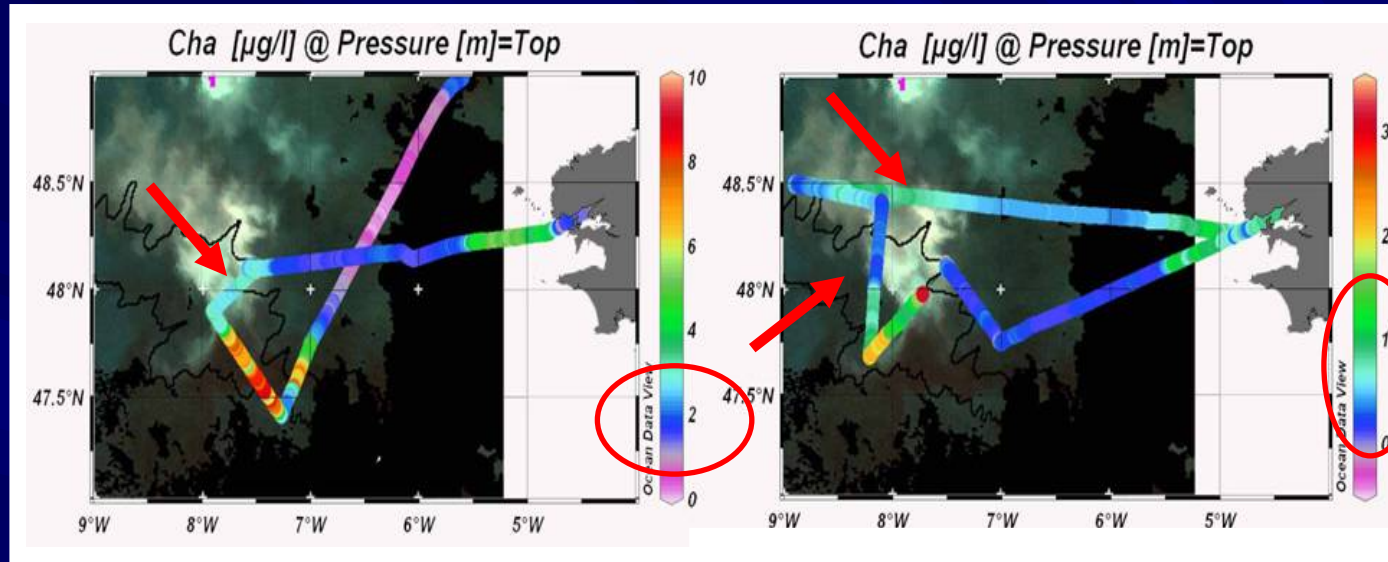
$[Chl-a] = 0.5 - 2 \mu g/l$



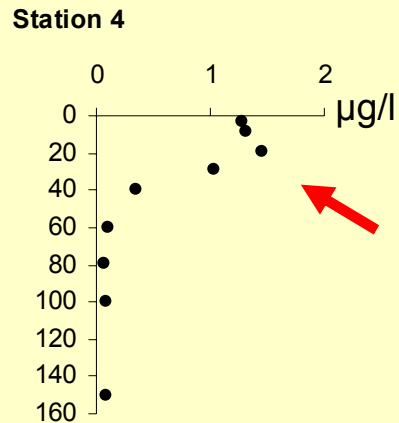
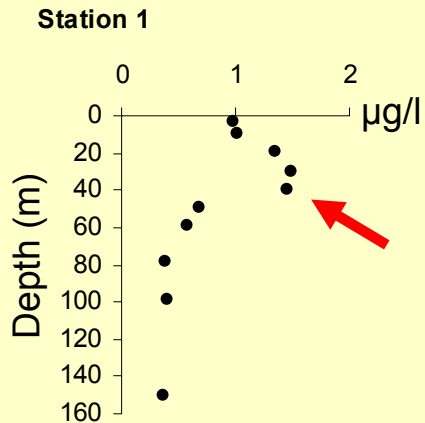


Chlorophyll-a concentrations

1st Leg



2nd Leg



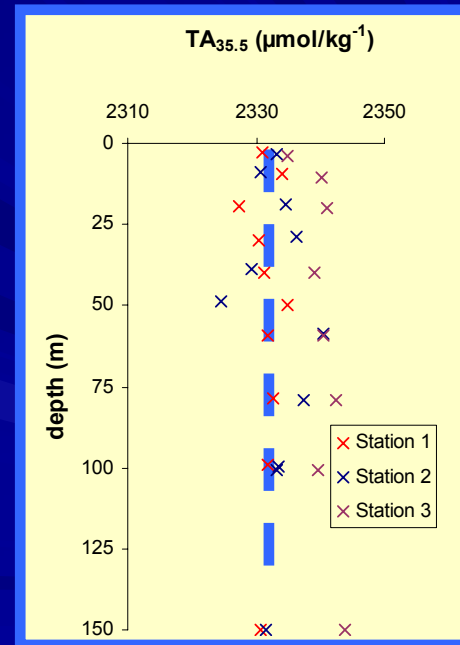
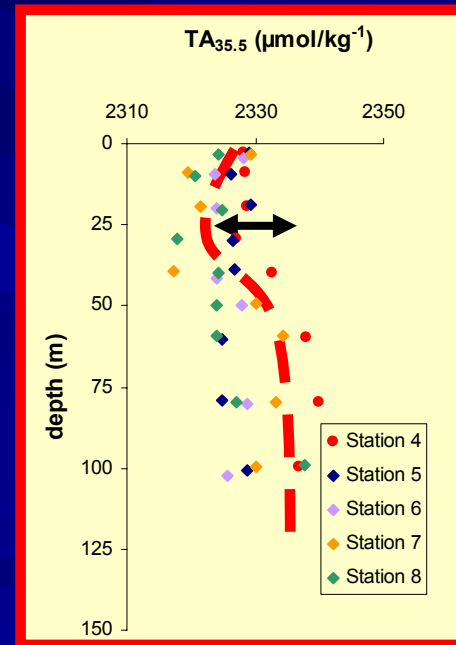
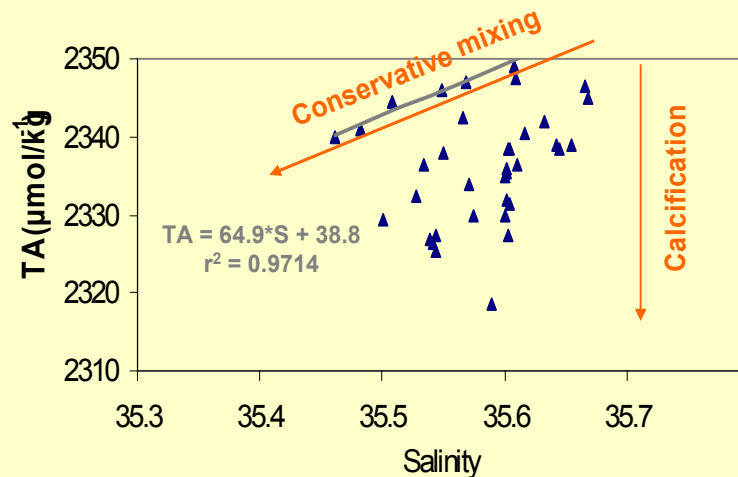
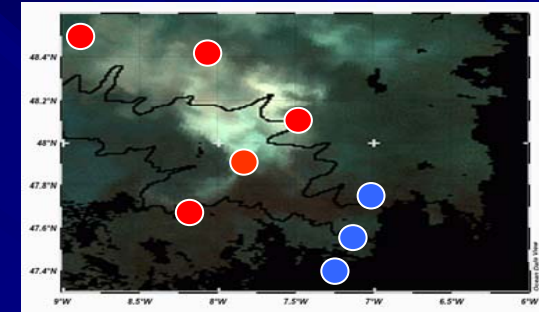
Moderate [Chl-a]

Deep Chl-a maximum ~ 20-40m



Total alkalinity

Biogenic calcification:

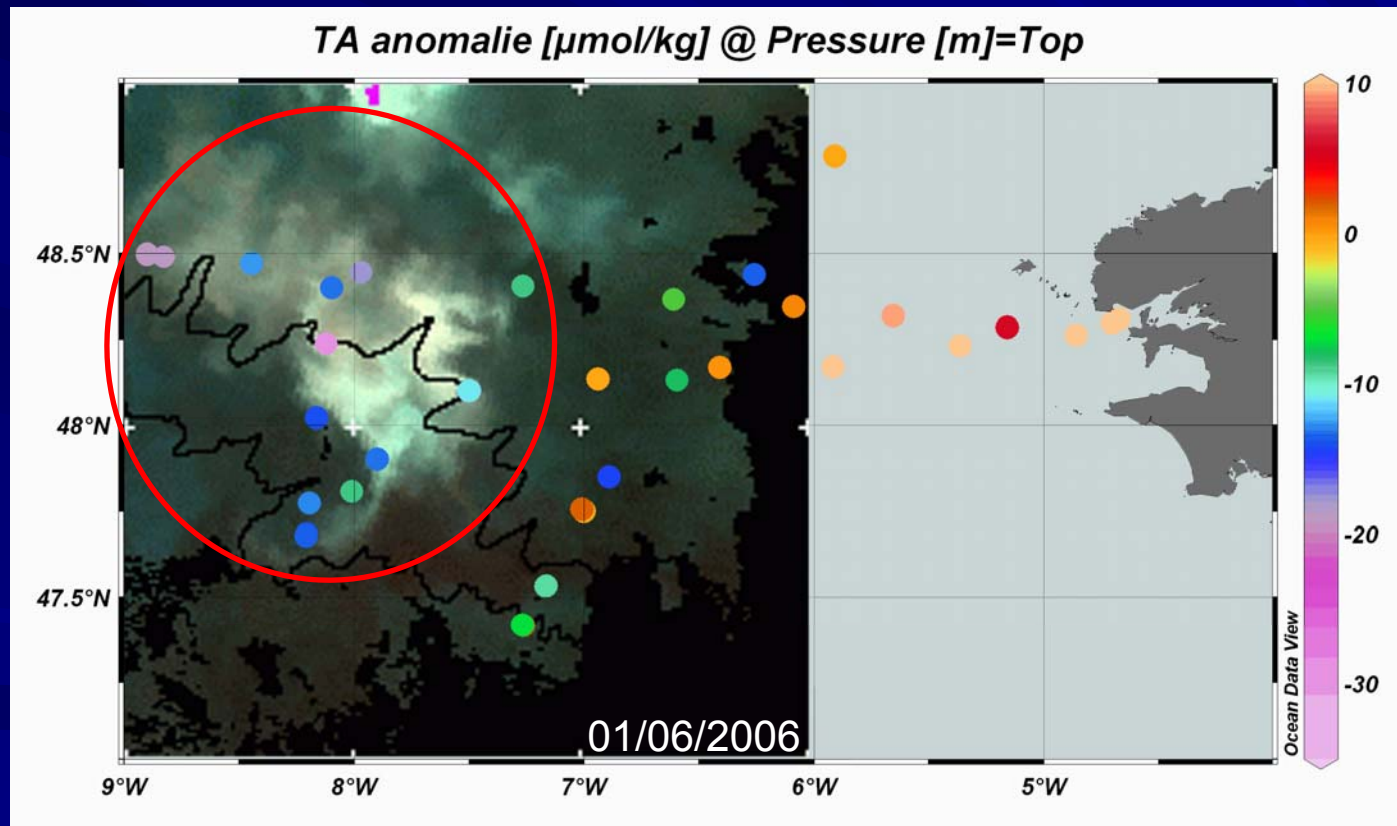


Fingerprint of calcification inside the
high reflectance patch



Total alkalinity anomalies

$$TA_{\text{anomaly}} = TA_{\text{meas}} - TA_{\text{sal}} \longrightarrow \text{maximum range of } 26 \mu\text{mol kg}^{-1}$$

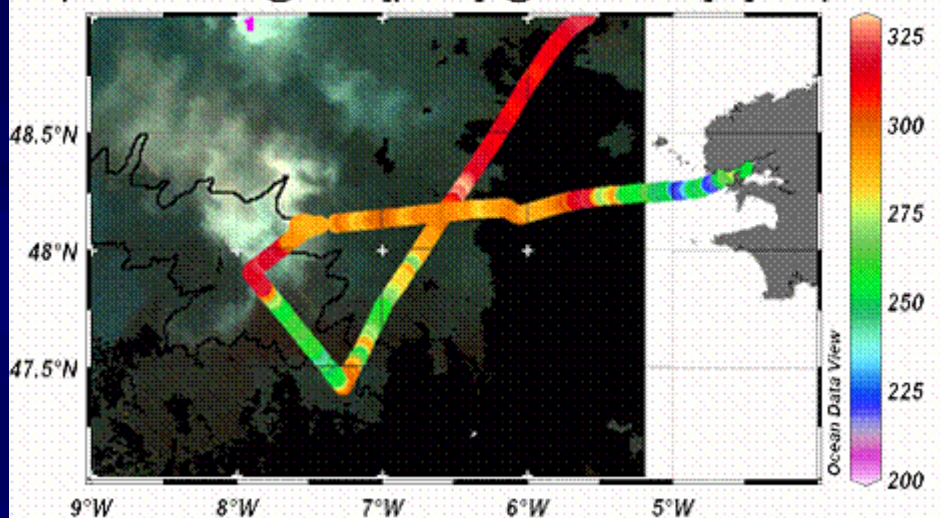


Details in Suykens *et al.* (**Poster BG0021 Today**)
Dissolved inorganic carbon dynamics in the Gulf of Biscay (June 2006)

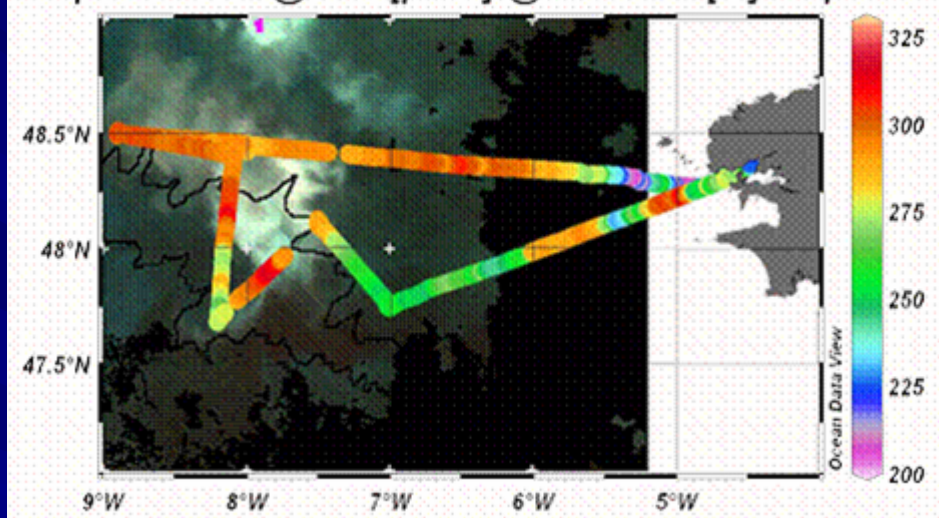


Sea-surface pCO₂

pCO₂ moist@13°C [μatm] @ Pressure [m]=Top

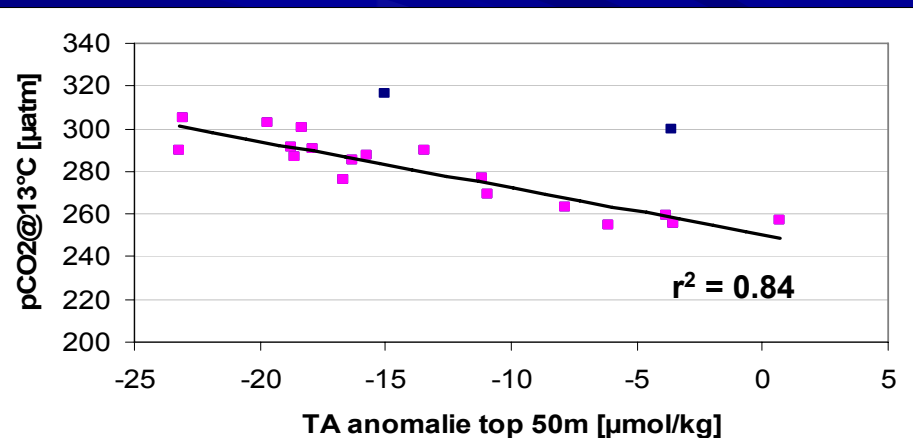


pCO₂ moist@13°C [μatm] @ Pressure [m]=Top



ranged from **250** to **338** μatm → sink for atmospheric CO₂ in early June 2006

strong correlation: TA anomaly vs. pCO₂@13°





Primary production and calcification

Primary production

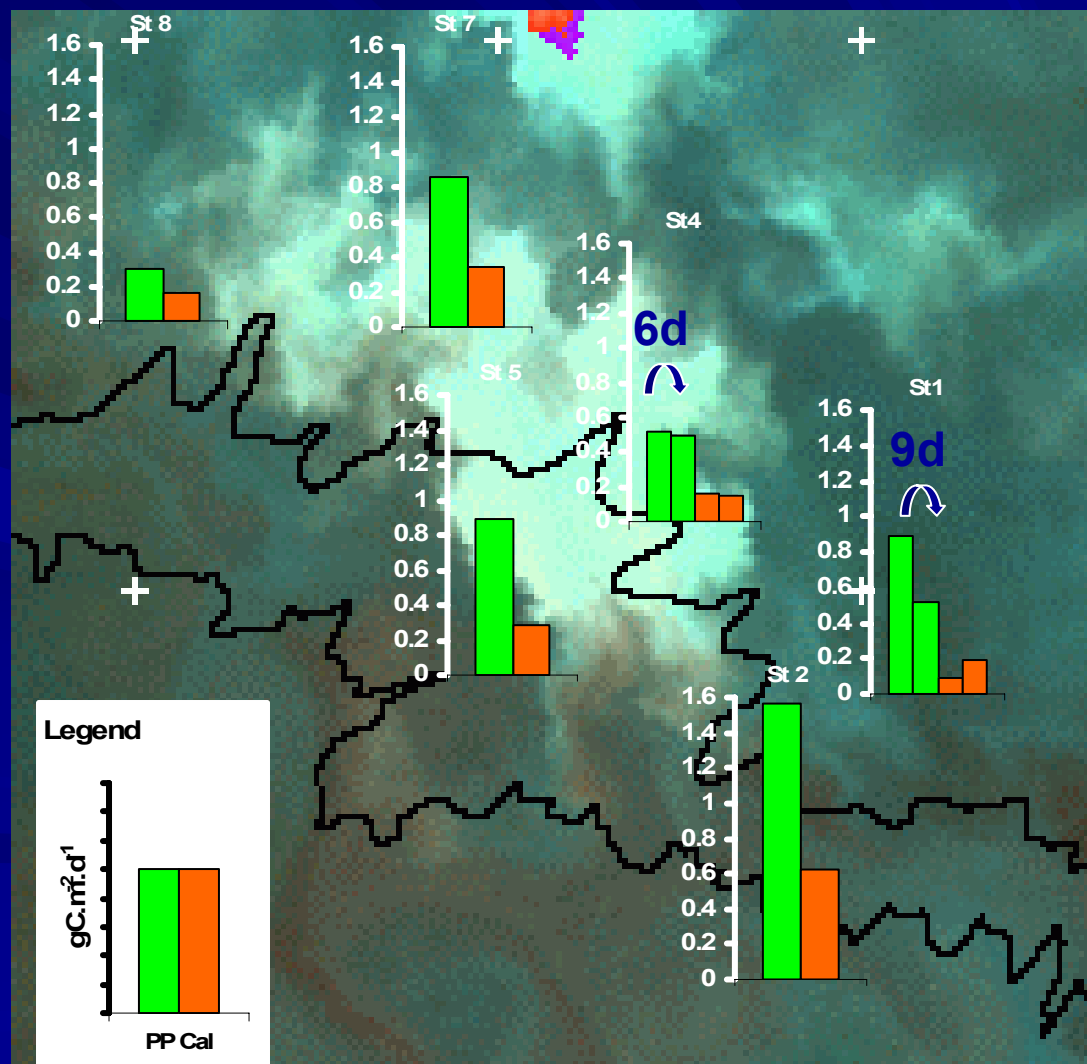
< 1 gC m⁻² d⁻¹ inside the patch
> 1.5 gC m⁻² d⁻¹ at station 2

Calcification rates

0.2 - 0.5 gC m⁻² d⁻¹

PIC:POC ~ 0.4
(molar uptake ratio)

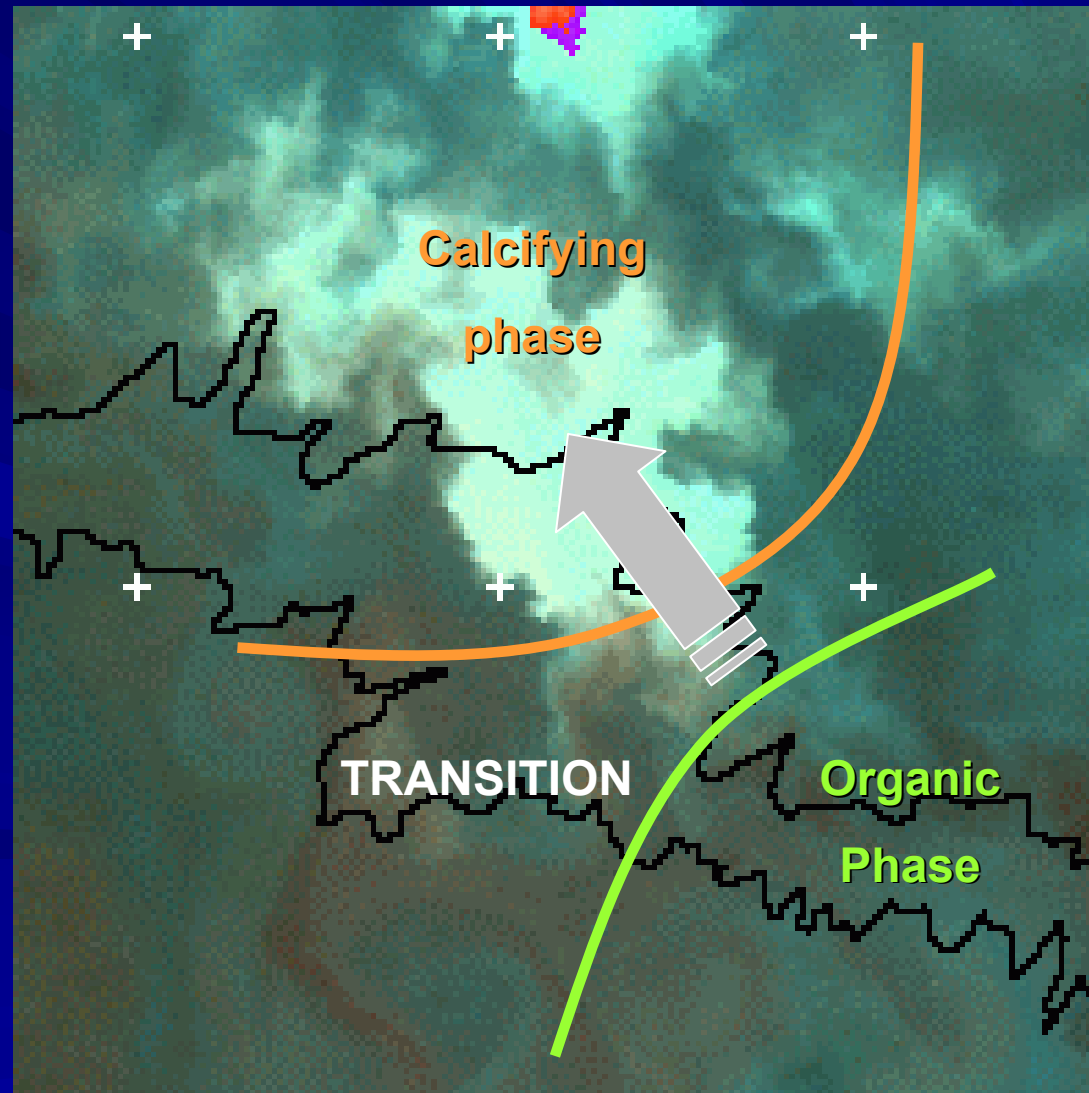
**Primary production decreased at
stations 1 and 4**





Primary production and calcification

- Zone dominated by the **organic phase** (1, 2)
- Zone of **transition** (4, 5)
- Zone dominated by the **calcifying phase** (7, 8)





Pelagic respiration rates

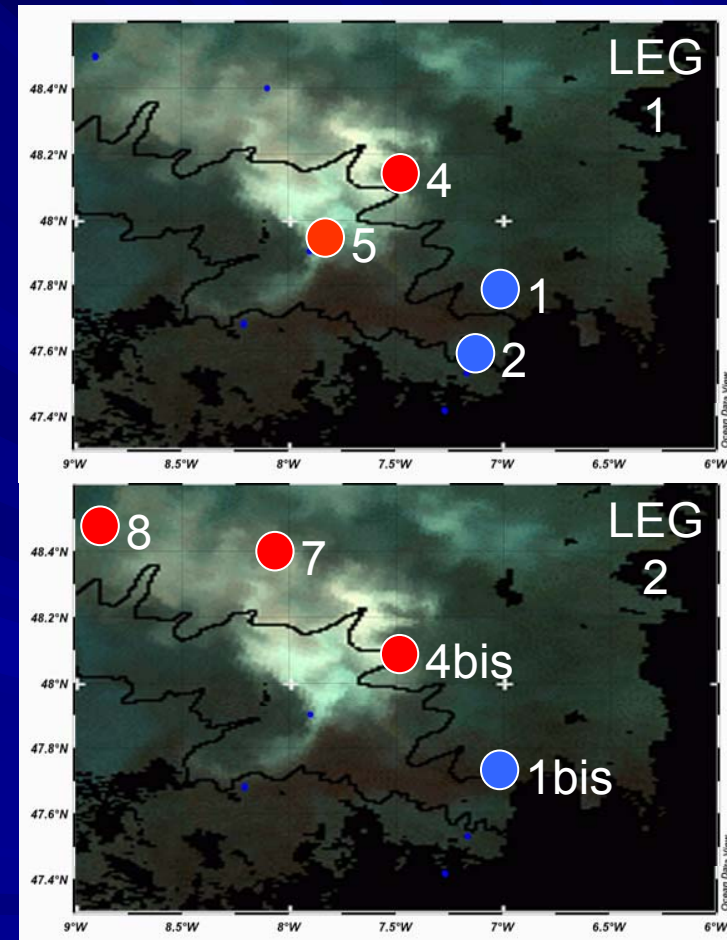
Station	PP	CAL	RESP (mmolC.m ⁻² .d ⁻¹)
1	74.2	7.5	82.3
1bis	43.3	15.8	116.9
2	130.8	51.7	90.7
5	74.2	24.2	
4	43.3	13.3	86.3
4bis	41.7	12.5	114.9
7	71.7	28.3	88.3
8	25.0	13.3	106.8

Ranging from 82 to 117 mmolC m⁻² d⁻¹

Total respiration as a major source of CO₂
(76 → 90 %)

Respiration rates exceed primary production rates,
except at st 2

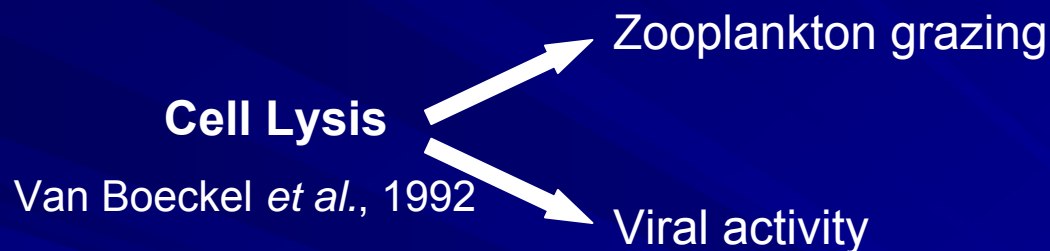
Increase at revisited stations





Lysis rate

Dissolved Esterase Activity method (Riegman *et al.*, 2002)

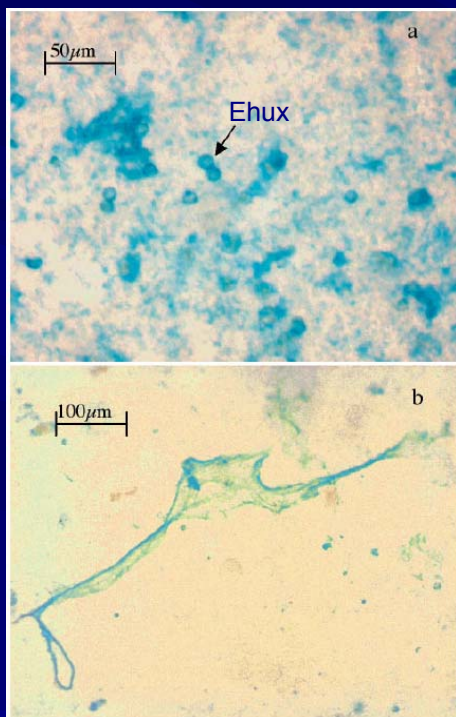


Station	date	Lysis (d ⁻¹)
2	1/06/2006	0.320
4	2/06/2006	0.253
8	6/06/2006	0.318
7	7/06/2006	0.178
1bis	9/06/2006	1.339

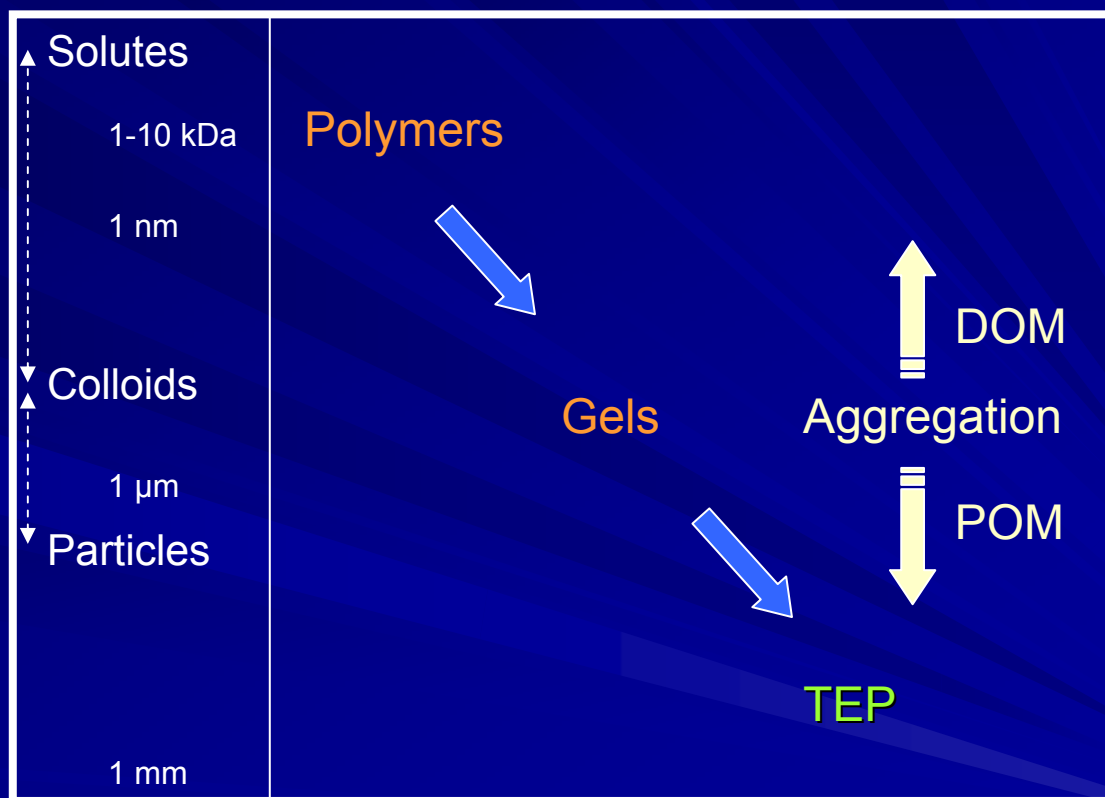
Typical for a late-spring bloom situation
0.02-0.3 d⁻¹ in Riegman & Winter (2003)



Transparent Exopolymer Particles



From Engel *et al.*, (2004)



Adapted from Verdugo *et al.*, (2004)

See also De Bodt *et al.* (Poster XY0734 **Wednesday**)

*Calcification and transparent exopolymer particles (TEP) production in batch cultures of *Emiliana huxleyi* exposed to different pCO₂*



Transparent Exopolymer Particles

Our study:

500-800 and up to **2000** $\mu\text{g Xeq/l}$

Engel (2004):

28 to 120 $\mu\text{g Xeq/l}$

NAO (June-July 1996)

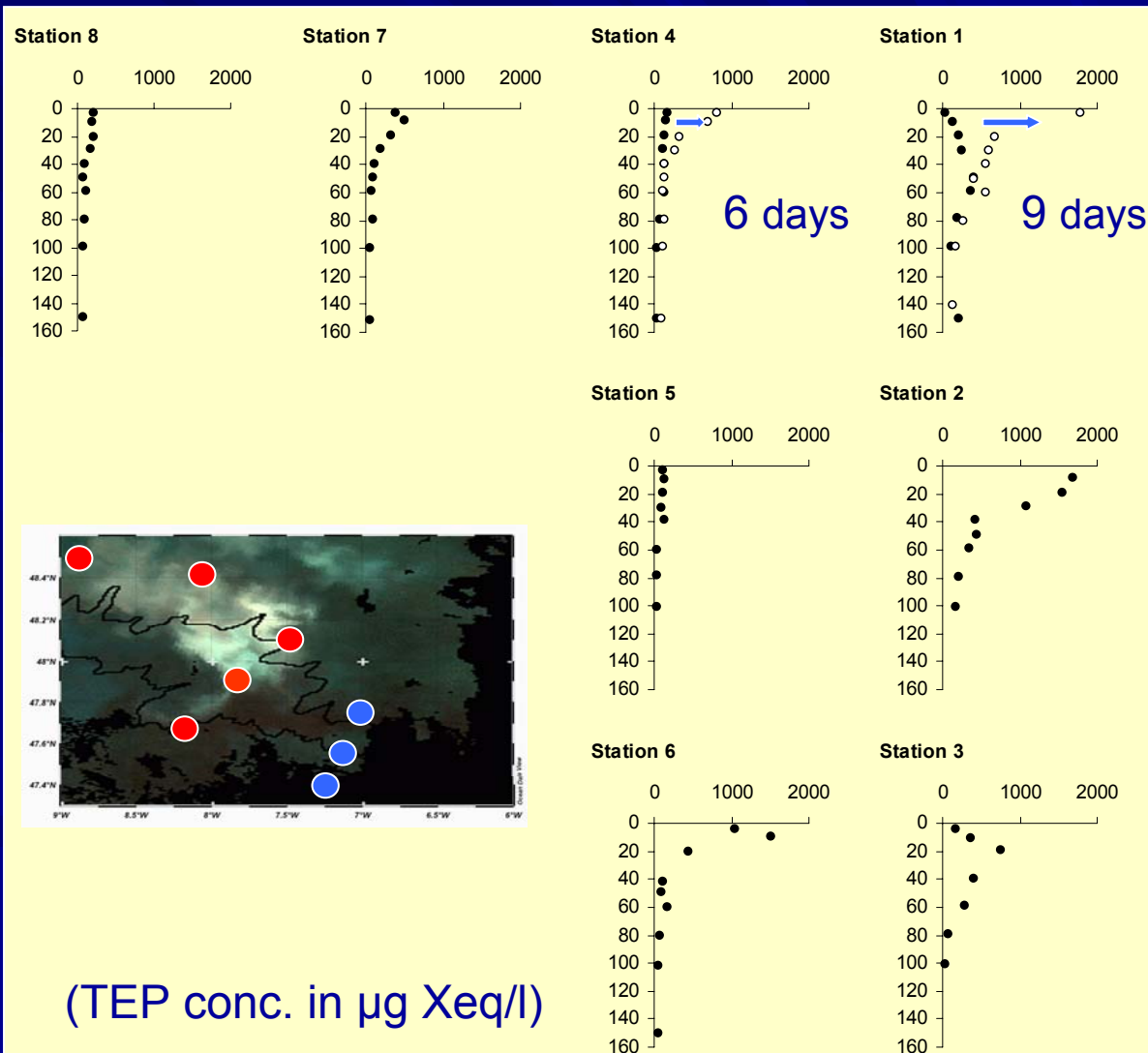
Passow *et al.* (2001):

500 $\mu\text{g Xeq/l}$ (Santa Barbara Channel)

Passow & Alldredge (1995):

920 $\mu\text{g Xeq/l}$ (batch culture, *Ehux*)

**Size spectrum and
specific polysaccharide
composition ?**





Bacterial community structure

DGGE / PCR 16s rDNA



« particle-associated fraction »

(> 3 μm)

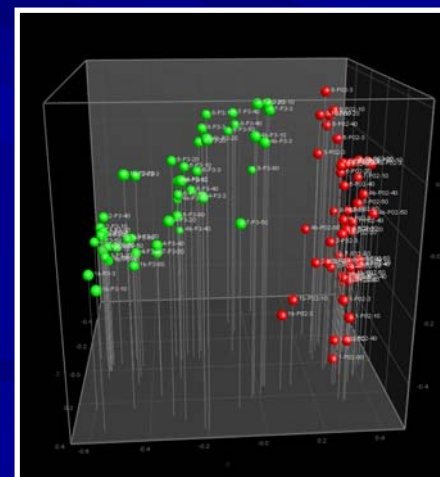
« free-living fraction »

(0.2 – 3 μm)

- α -proteobacteria (*Roseobacter*)
- SAR86 lineage



DMS-cleavage pathway
Organic-S metabolism



- Particle-associated fraction
- Free-living fraction



CONCLUSIONS

The continental margin has hydrodynamic features that enhance biological activity and especially promote **coccolithophore blooms**.

This ecosystem was still a **sink for atmospheric CO₂** due to the history of the bloom development.

But the intensity of the CO₂-producing processes is important and may switch the system from a sink to a source of CO₂.

The **elevated concentrations of TEP**, accompanied by **high cell lysis rates**, may lead to the **production and subsequent export of macro-aggregates**, which could be enhanced by the ballasting of calcite particles.

“What is the role of bacteria in aggregates ? Dissolution of CaCO₃ ? DMS-cleavage pathway ?”

Contribution to C export and preservation via shelf-ocean exchanges

Future of shelf carbonate sediments in a high-CO₂ world...



Thank you for your attention

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